

IFW16

RAW SEQUENCE LISTING DATE: 07/29/2004
PATENT APPLICATION: US/09/618,380C TIME: 09:31:19

Input Set : A:\SEQLIST4920USv3.txt

Output Set: N:\CRF4\07292004\I618380C.raw

```
3 <110> APPLICANT: Weiner, George
 4 Gingrich, Roger
       Link, Brian
       Tso, J. Yun
8 <120> TITLE OF INVENTION: HUMANIZED ANTIBODIES AGAINST CD3
10 <130> FILE REFERENCE: 05882-0176-CNUS03
12 <140> CURRENT APPLICATION NUMBER: US 09/618,380C
13 <141> CURRENT FILING DATE: 2000-07-18
15 <150> PRIOR APPLICATION NUMBER: US 08/397,411
16 <151> PRIOR FILING DATE: 1995-03-01
18 <150> PRIOR APPLICATION NUMBER: US 07/859,583
19 <151> PRIOR FILING DATE: 1992-03-27
21 <160> NUMBER OF SEQ ID NOS: 14
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23 <170> SOFTWARE: PatentIn version 3.1
25 <210> SEQ ID NO: 1
26 <211> LENGTH: 107
27 <212> TYPE: PRT
28 <213> ORGANISM: Artificial Sequence
30 <220> FEATURE:
31 <223> OTHER INFORMATION: Light chain of Humanized 1D10 Ab minus signal sequence
33 <400> SEQUENCE: 1
35 Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
39 Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Glu Asn Ile Tyr Ser Tyr
40 20
                                 25
43 Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Val
                             40
47 Ser Asn Ala Lys Thr Leu Ala Glu Gly Val Pro Ser Arg Phe Ser Gly
                         55
51 Ser Gly Ser Gly Lys Gln Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro
55 Glu Asp Phe Ala Thr Tyr Tyr Cys Gln His His Tyr Gly Asn Ser Tyr
59 Pro Phe Gly Gln Gly Thr Lys Leu Glu Ile Lys
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63 <210> SEQ ID NO: 2
64 <211> LENGTH: 107
65 <212> TYPE: PRT
66 <213> ORGANISM: Mus sp.
68 <400> SEQUENCE: 2
70 Asp Ile Gln Met Thr Gln Ser Pro Ala Ser Leu Ser Ala Ser Val Gly
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74 Glu Thr Val Thr Ile Thr Cys Arg Ala Ser Glu Asn Ile Tyr Ser Tyr

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```
78 Leu Ala Trp Tyr Gln Gln Lys Gln Gly Lys Ser Pro Gln Leu Leu Val
           35
                               40
82 Ser Asn Ala Lys Thr Leu Ala Glu Gly Val Thr Ser Arg Phe Ser Gly
                           55
86 Ser Gly Ser Gly Lys Gln Phe Ser Leu Lys Ile Asn Ser Leu Gln Pro
                                            75
90 Glu Asp Phe Gly Asn Tyr Tyr Cys Gln His His Tyr Gly Asn Ser Tyr
94 Pro Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys
95
               100
98 <210> SEQ ID NO: 3
99 <211> LENGTH: 116
100 <212> TYPE: PRT
101 <213> ORGANISM: Artificial Sequence
103 <220> FEATURE:
104 <223> OTHER INFORMATION: Heavy chain of Humanized 1D10 Ab minus signal sequence
106 <400> SEQUENCE: 3
108 Gln Val Gln Leu Gln Glu Ser Gly Pro Gly Leu Val Lys Pro Ser Glu
109 1
                                        1.0
112 Thr Leu Ser Leu Thr Cys Thr Val Ser Gly Phe Ser Leu Thr Asn Tyr
116 Gly Val His Trp Val Arg Gln Ser Pro Gly Lys Gly Leu Glu Trp Ile
                                                           Sec. 1
                                                     45
120 Gly Val Lys Trp Ser Gly Gly Ser Thr Glu Tyr Asn Ala Ala Phe ile
124 Ser Arg Leu Thr Ile Ser Lys Asp Thr Ser Lys Asn Gln Val Ser Leu
125 65
                        70
                                            75
128 Lys Leu Asn Ser Leu Thr Ala Ala Asp Thr Ala Val Tyr Tyr Cys Ala
                    85
                                        90
132 Arg Asn Asp Arg Tyr Ala Met Asp Tyr Trp Gly Gln Gly Thr Leu Val
                100
                                    105
136 Thr Val Ser Ser
137
            115
140 <210> SEQ ID NO: 4
141 <211> LENGTH: 116
142 <212> TYPE: PRT
143 <213> ORGANISM: Mus sp.
145 <400> SEQUENCE: 4
147 Gln Val Gln Leu Lys Gln Ser Gly Pro Gly Leu Val Gln Pro Ser Gln
151 Ser Leu Ser Ile Thr Cys Thr Gly Ser Gly Phe Ser Leu Thr Asn Tyr
155 Gly Val His Trp Val Arg Gln Ser Pro Gly Lys Gly Leu Glu Trp Leu
159 Gly Val Lys Trp Ser Gly Gly Ser Thr Glu Tyr Asn Ala Ala Phe Ile
163 Ser Arg Leu Ser Ile Ser Lys Asp Asn Ser Lys Ser Gln Val Phe Phe
164 65
                        70
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```
167 Lys Met Asn Ser Leu Gln Ala Asp Asp Thr Ala Met Tyr Tyr Cys Ala
168
171 Arg Asn Asp Arg Tyr Ala Met Asp Tyr Trp Gly Gln Gly Thr Ser Val
                100
172
                                     105
175 Thr Val Ser Ser
176
            115
179 <210> SEQ ID NO: 5
180 <211> LENGTH: 214
181 <212> TYPE: PRT
182 <213> ORGANISM: Artificial Sequence
184 <220> FEATURE:
185 <223> OTHER INFORMATION: Complete light chain of Humanized 1D10 Ab
187 <400> SEQUENCE: 5
189 Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
193 Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Glu Asn Ile Tyr Ser Tyr
                20
197 Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Val
198
            35
                                 40
201 Ser Asn Ala Lys Thr Leu Ala Glu Gly Val Pro Ser Arq Phe Ser Gly
                            55
205 Ser Gly Ser Gly Lys Gln Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro
209 Glu Asp Phe Ala Thr Tyr Tyr Cys Gln His His Tyr Gly Asn Ser Tyr
213 Pro Phe Gly Gln Gly Thr Lys Leu Glu Ile Lys Arg Thr Val Ala Ala
                                     105
217 Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu Gln Leu Lys Ser Gly
218
                                 120
            115
                                                     125
221 Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe Tyr Pro Arg Glu Ala
                            135
                                                 140
225 Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln Ser Gly Asn Ser Gln
                        150
                                             155
229 Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser Thr Tyr Ser Leu Ser
                                         170
233 Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu Lys His Lys Val Tyr
234
                180
                                    185
237 Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser Pro Val Thr Lys Ser
            195
                                200
241 Phe Asn Arg Gly Glu Cys
242
        210
245 <210> SEQ ID NO: 6
246 <211> LENGTH: 273
247 <212> TYPE: PRT
248 <213> ORGANISM: Artificial Sequence
250 <220> FEATURE:
251 <223> OTHER INFORMATION: Fd-jun in F(ab'-zipper)2 of humanized 1D10 antibody
253 <400> SEQUENCE: 6
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255 Gln Val Gln Leu Gln Glu Ser Gly Pro Gly Leu Val Lys Pro Ser Glu

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Input Set : A:\SEQLIST4920USv3.txt

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256	1				5					10					15		
259	Thr	Leu	Ser	Leu	Thr	Cys	Thr	Val	Ser	Gly	Phe	Ser	Leu	Thr	Asn	Tyr	
260				20		-			25	_				30		-	
263	Gly	Val	His	Trp	Val	Arg	Gln	Ser	Pro	Gly	Lys	Gly	Leu	Glu	Trp	Ile	
264			35			•		40					45				
267	Gly	Val	Lys	Trp	Ser	Gly	Gly	Ser	Thr	Glu	Tyr	Asn	Ala	Ala	Phe	Ile	
268		50	•				55					60					
271	Ser	Arg	Leu	Thr	Ile	Ser	Lys	Asp	Thr	Ser	Lys	Asn	Gln	Val	Ser	Leu	
272	65					70					75					80	
275	Lys	Leu	Asn	Ser	Leu	Thr	Ala	Ala	Asp	Thr	Ala	Val	Tyr	Tyr	Cys	Ala	
276					85					90					95		
279	Arg	Asn	Asp	Arg	Tyr	Ala	Met	Asp	Tyr	Trp	Gly	Gln	Gly	Thr	Leu	Val	
280				100		•			105					110			
283	Thr	Val	Ser	Ser	Ala	Ser	Thr	Lys	Gly	Pro	Ser	Val	Phe	Pro	Leu	Ala	
28 4			115					120					125				
2 87	Pro	Ser	Ser	Lys	Ser	Thr	Ser	Gly	Gly	Thr	Ala	Ala	Leu	Gly	Cys	Leu	
288		130					135					140					
291	Val	Lys	Asp	Tyr	Phe	Pro	Glu	Pro	Val	Thr	Val	Ser	Trp	Asn	Ser	Gly	
292	145					150					155					160	
295	Ala	Leu	Thr	Ser	Gly	Val	His	Thr	Phe	Pro	Ala	·Val	Leu	Gln	Ser	Ser	
296					165					170					175		
299	Gly	Leu	Tyr	Ser	Leu	Ser	Ser	Val	Val	Thr	Val	Pro	Ser	Ser	Ser	Leu	
300				180					185					190			
303	Gly	Thr	Gln	Thr	Tyr	Ile	Cys	Asn	Val	Asn	His	Lys	Pro	Ser	Asn	Thr	
304			195					200					205				
307	Lys	Val	Asp	Lys	Lys	Val	Glu	Pro	Lys	Ser	Cys	Asp	Lys	Thr	His	Thr	
308		210					215					220					
311	Cys	Pro	Pro	Cys	Lys	Cys	Pro	Ala	Gly	Gly		Ile	Ala	Arg	Leu	Glu	
	225					230					235					240	
315	Glu	Lys	Val	Lys		Leu	Lys	Ala	Gln		Ser	Glu	Leu	Ala	Ser	Thr	
316	_				245	_	_		_	250					255		
	Ala	Asn	Met		Arg	Glu	Gln			Gln	Leu	Lys	Gln		Val	Met	
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	Asn				_												
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	<pre><212> TYPE: PRT </pre> <pre><213> ORGANISM: Artificial Sequence</pre>																
					Art:	LT1C:	ial S	Seque	ence								
	2 <220> FEATURE:																
	3 <223> OTHER INFORMATION: Complete heavy chain of Humanized 1D10 Ab																
			~			~ 7	_	~ 1	_	~-7	_		_	_	_	~7	
		Val	Gln	ьeu		Glu	Ser	GIY	Pro	_	Leu	Val	ьуs	Pro		Glu	
338		_	~	-	5	~	m)		<u> </u>	10	D1			m\	15		
	Tnr	ьeu	Ser		Inr	Cys	Tnr	val		GTA	Pne	ser	ьeu		Asn	Tyr	
342	0.7	YY . 3	***	20	17. 3	20	0.7	0	25	a 3 -	T	a a	T	30	m-···	T7 .	
	GIA	val	His	Trp	vaı	arg	GIN		Pro	GIY	ьys	GTA		GIU	Trp	ше	
346	G 7	77. 7	35		a .	a 7	a 3	40	ml	~ 3		70 .	45	7. 7	D1	- 7	
	GTÀ		Lys	rrp	ser	GIÀ		ser	ınr	GIU	Tyr		Ala	АТА	rne	тте	
350		50					55					60	-				

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353 354		Arg	Leu	Thr	Ile	Ser 70	Lys	Asp	Thr	Ser	Lys 75	Asn	Gln	Val	Ser	Leu 80
357 358	Lys	Leu	Asn	Ser	Leu 85	Thr	Ala	Ala	Asp	.Thr 90	Ala	Val	Tyr	Tyr	Cys 95	Ala
	Arg	Asn	Asp	Arg		Ala	Met	Asp	Tyr 105		Gly	Gln	Gly	Thr 110		Val
	Thr	Val	Ser 115	Ser	Ala	Ser	Thr	Lys 120	Gly	Pro	Ser	Val	Phe 125		Leu	Ala
369 370	Pro	Ser 130	Ser	Lys	Ser	Thr	Ser 135	Gly	Gly	Thr	Ala	Ala 140	Leu	Gly	Cys	Leu
	Val 145	Lys	Asp	Tyr	Phe	Pro 150	Glu	Pro	Val	Thr	Val 155	Ser	Trp	Asn	Ser	Gly 160
		Leu	Thr	Ser	Gly 165		His	Thr	Phe	Pro 170		Val	Leu	Gln	Ser 175	
	Gly	Leu	Tyr	Ser 180		Ser	Ser	Val	Val 185		Val	Pro	Ser	Ser 190		Leu
	Gly	Thr	Gln 195	Thr	Tyr	Ile	Суѕ	Asn 200	Val	·Asn	His	Lys	Pro 205	Ser	Asn	Thr
	Lys	Val 210	Asp	Lys	Lys	Val	Glu 215		Lys	Ser	Сув	Asp 220	Lys	Thr	His	Thr
	Cys 225	Pro	Pro	Cys	Pro	Ala 230	Pro	Glu	Leu	Leu	Gly 235	Gly	Pro	Ser	Val	Phe 240
		Phe	Pro	Pro	Lys 245		Lys	Asp	Thr	Leu 250		Ile	Ser	Arg	Thr 255	Pro
401 402	Glu	Val	Thr	Cys 260	Val	Val	Val	Asp	Val 265	Ser	His	Glu	Asp	Pro 270	Glu	Val
405 406	Lys	Phe	Asn 275	Trp	Tyr	Val	Asp	Gly 280	Val	Glu	Val	His	Asn 285	Ala	Lys	Thr
409 410	Lys	Pro 290	Arg	Glu	Glu	Gln	Tyr 295	Asn	Ser	Thr	Tyr	Arg 300	Val	Val	Ser	Val
	Leu 305	Thr	Val	Leu	His	Gln 310	Asp	Trp	Leu	Asn	Gly 315	Lys	Glu	Tyr	Lys	Cys 320
417 418	Lys	Val	Ser	Asn	Lys 325	Ala	Leu	Pro	Ala	Pro 330	Ile	Glu	Lys	Thr	Ile 335	Ser
421 422	Lys	Ala	Lys	Gly 340	Gln	Pro	Arg	Glu	Pro 345	Gln	Val	Tyr	Thr	Leu 350	Pro	Pro
425 426	Ser	Arg	Asp 355	Glu	Leu	Thr	Lys	Asn 360	Gln	Val	Ser	Leu	Thr 365	Cys	Leu	Val
429 430	Lys	Gly 370	Phe	Tyr	Pro	Ser	Asp 375	Ile	Ala	Val	Glu	Trp 380	Glu	Ser	Asn	Gly
	Gln 385	Pro	Glu	Asn	Asn	Tyr 390	Lys	Thr	Thr	Pro	Pro 395	Val	Leu	Asp	Ser	Asp
437 438	Gly	Ser	Phe	Phe	Leu 405	Tyr	Ser	Lys	Leu	Thr 410	Val	Asp	Lys	Ser	Arg 415	Trp
441 442	Gln	Gln	Gly	Asn 420	Val	Phe	Ser	Cys	Ser 425	Val	Met	His	Glu	Ala 430	Leu	His
	Asn	His	Tyr 435		Gln	Lys	Ser	Leu 440		Leu	Ser	Pro	Gly 445			
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RAW SEQUENCE LISTING ERROR SUMMARY

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:14; Xaa Pos. 2,3,4,5,6,7

VERIFICATION SUMMARY

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L:767 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:0